#### **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/581.354
Source:	IFWP.
Date Processed by STIC:	6/19/06
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## ENTERED

### CRF Errors Edited by the STIC Systems Branch

Serial	Number: <u>/0/58/, 354</u>	CRF Edit Date: 6/19/0
· 	Realigned nucleic acid/amino acid numbers/text text "wrapped" to the next line	in cases where the sequence
	Corrected the SEQ ID NO. Sequence numbers e	edited were:
· 	Inserted or corrected a nucleic number at the en NO's edited:	d of a nucleic line. SEQ ID
_	Deleted: invalid beginning/end-of-file text;	page numbers
	Inserted mandatory headings/numeric identifier	s, specifically:
	Moved responses to same line as heading/numer	ic identifier, specifically:
	Other:	

Revised 09/09/2003



IFWP

RAW SEQUENCE LISTING DATE: 06/19/2006
PATENT APPLICATION: US/10/581,354 TIME: 11:38:20

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3 <110> APPLICANT: INSTITUT PASTEUR

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              CRESCENZO-CHAIGNE, Bernadette
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             GERBAUD, Sylvie
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             BURGUIERE, Ana Maria
             AZEBI, Saliha
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             CHARNEAU, Pierre
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             DELAGNEAU, Jean-Francois
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             MARTIN, Monique
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STRAIN OF
              SARS-ASSOCIATED CORONAVIRUS
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C--> 28 <141> CURRENT FILING DATE: 2006-06-02
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VERIFICATION SUMMARY DATE: 06/19/2006
PATENT APPLICATION: US/10/581,354 TIME: 11:38:21

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06192006\J581354.raw

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8 CRESCENZO-CHAIGNE, Bernadette
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10 KUNST, Franck
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STRAIN OF
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C--> 28 <140> CURRENT APPLICATION NUMBER: US/10/581,354
C--> 28 <141> CURRENT FILING DATE: 2006-06-02
      28 <150> PRIOR APPLICATION NUMBER: FR 0314152
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      36 <170> SOFTWARE: PatentIn version 3.1
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#### **ERRORED SEQUENCES**

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E--> 5257
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/581,354

DATE: 06/14/2006
TIME: 10:08:36

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\06142006\J581354.raw

L:28 M:270 C: Current Application Number differs, Replaced Current Application No L:28 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:551 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:549 L:1189 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:9,Line#:1187 L:1302 M:283 W: Missing Blank Line separator, <400> field identifier L:1302 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:1301 L:1382 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:1380 L:1443 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:1441 L:1609 M:283 W: Missing Blank Line separator, <400> field identifier L:1609 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:21,Line#:1608 L:1670 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:23,Line#:1668 L:1746 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:25, Line#:1744 L:1801 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:27, Line#:1799 L:1854 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:29, Line#:1852 L:2341 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:36, Line#:2339 L:4213 M:283 W: Missing Blank Line separator, <400> field identifier L:4337 M:283 W: Missing Blank Line separator, <400> field identifier L:4391 M:283 W: Missing Blank Line separator, <400> field identifier L:4522 M:283 W: Missing Blank Line separator, <400> field identifier L:4883 M:283 W: Missing Blank Line separator, <400> field identifier L:5077 M:283 W: Missing Blank Line separator, <400> field identifier L:5257 M:254 E: No. of Bases conflict, this line has no nucleotides.